```
===============================================================
Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: markspencer
Timestamp: [year=2008; month=1; day=11; hr=17; min=2; sec=10; ms=804; ]
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*************************************************************
Reviewer Comments:
Please realign amino acid number so that it lines up with every fifth
amino acid.
*************************************************************
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Application No: 10572522   Version No: 1.0

Input Set:

Output Set:

Started:	2007-12-18 14:57:25.234
Finished:	2007-12-18 14:57:26.071
Elapsed:	0 hr(s) 0 min(s) 0 sec(s) 837 ms
Total Warnings:	0
Total Errors:	5
No. of SeqIDs Defined:	1
Actual SeqID Count:	1

Error code	Error Description
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (20)
E 323	Invalid/missing amino acid numbering SEQID (1) POS (24)
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (25)
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E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (30)

SEQUENCE LISTING

<110>   Andreas BERGMANN

<120>   METHOD FOR THE DIAGNOSIS OF DISEASES BY DETERMINING APOLIPOPROTEIN C-1

<130>   2582.011

<140>   10572522
<141>   2007-12-18

<150>   PCT/EP2004/09965
<151>   2004-09-07

<150>   DE 103 43 815.7
<151>   2003-09-22

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<213>   Homo sapiens

<400>   1

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                      20   25   30

Glu Leu Ser Ala Lys Met Arg Glu Trp Phe Ser Glu Thr Phe Gln Lys
        35   40   45

Val Lys Glu Lys Leu Lys Ile Asp Ser
    50   55